

## SEQUENCE LISTING

<110> THE JOHNS HOPKINS UNIVERSITY  
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<120> METHODS OF IDENTIFYING MODULATORS OF CELLULAR GLYCOSYLATION USING  
 GTRAP3-18

<130> JHU2090-1

<140> US 10/542,435  
 <141> 2004-01-18

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 <151> 2004-01-18

<150> US 60/440,717  
 <151> 2003-01-17

<160> 10

<170> PatentIn version 3.3

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 <212> DNA  
 <213> Rattus norvegicus

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 ctctattacc agaccaacta cctggtgggtg gctgccatga tgatttcagt cgttggggttt 180  
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 cttcgaaacc tcaagaacaa actggaaaat aaaatggagg gaataggctt gaagaaaacg 480  
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Gly Ser Asp Arg Phe Ala Arg Pro Asp Phe Arg Asp Ile Ser Lys Trp  
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Asn Asn Arg Val Val Ser Asn Leu Leu Tyr Tyr Gln Thr Asn Tyr Leu  
 35 40 45

Val Val Ala Ala Met Met Ile Ser Val Val Gly Phe Leu Ser Pro Phe  
 50 55 60

Asn Met Ile Leu Gly Gly Ile Ile Val Val Leu Val Phe Thr Gly Phe  
 65 70 75 80

Val Trp Ala Ala His Asn Lys Asp Ile Leu Arg Arg Met Lys Lys Gln  
 85 90 95

Tyr Pro Thr Ala Phe Val Met Val Val Met Leu Ala Ser Tyr Phe Leu  
 100 105 110

Ile Ser Met Phe Gly Gly Val Met Val Phe Val Phe Gly Ile Thr Phe  
 115 120 125

Pro Leu Leu Leu Met Phe Ile His Ala Ser Leu Arg Leu Arg Asn Leu  
 130 135 140

Lys Asn Lys Leu Glu Asn Lys Met Glu Gly Ile Gly Leu Lys Lys Thr  
 145 150 155 160

Pro Met Gly Ile Ile Leu Asp Ala Leu Glu Gln Gln Glu Asp Ser Ile  
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Asn Lys Phe Ala Asp Tyr Ile Ser Lys Ala Arg Glu  
 180 185

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<211> 96

<212> PRT

<213> Homo sapiens P429-End EAAT3

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Pro Ala Glu Asp Val Thr Leu Ile Ile Ala Val Asp Trp Leu Leu Asp  
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Arg Phe Arg Thr Met Val Asn Val Leu Gly Asp Ala Phe Gly Thr Gly  
 20 25 30

Ile Val Glu Lys Leu Ser Lys Lys Glu Leu Glu Gln Met Asp Val Ser

35                                      40                                      45  
 Ser Glu Val Asn Ile Val Asn Pro Phe Ala Leu Glu Ser Thr Ile Leu  
   50                                      55                                      60  
 Asp Asn Glu Asp Ser Asp Thr Lys Lys Ser Tyr Val Asn Gly Gly Phe  
  65                                      70                                      75                                      80  
 Ala Val Asp Lys Ser Asp Thr Ile Ser Phe Thr Gln Thr Ser Gln Phe  
                                     85                                      90                                      95  
  
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 Arg Phe Arg Thr Met Ile Asn Val Leu Gly Asp Ala Leu Ala Ala Gly  
                                     20                                      25                                      30  
 Ile Met Ala His Ile Cys Arg Lys Asp Phe Ala Arg Asp Thr Gly Thr  
                                     35                                      40                                      45  
 Glu Lys Leu Leu Pro Cys Glu Thr Lys Pro Val Ser Leu Gln Glu Ile  
  50                                      55                                      60  
 Val Ala Ala Gln Gln Asn Gly Cys Val Lys Ser Val Ala Glu Ala Ser  
  65                                      70                                      75                                      80  
 Glu Leu Thr Leu Gly Pro Thr Cys Pro His His Val Pro Val Gln Val  
                                     85                                      90                                      95  
 Glu Arg Asp Glu Glu Leu Pro Ala Ala Ser Leu Asn His Cys Thr Ile  
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 Gln Ile Ser Glu Leu Glu Thr Asn Val  
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Pro Thr His Asp Leu Pro Leu Ile Leu Ala Val Asp Trp Ile Val Asp  
 1 5 10 15

Arg Thr Thr Thr Val Val Asn Val Glu Gly Asp Ala Leu Gly Ala Gly  
 20 25 30

Ile Leu His His Leu Asn Gln Lys Ala Thr Lys Lys Gly Glu Gln Glu  
 35 40 45

Leu Ala Glu Val Lys Val Glu Ala Ile Pro Asn Cys Lys Ser Glu Glu  
 50 55 60

Glu Thr Ser Pro Leu Val Thr His Gln Asn Pro Ala Gly Pro Val Ala  
 65 70 75 80

Ser Ala Pro Glu Leu Glu Ser Lys Glu Ser Val Leu  
 85 90

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 <211> 93  
 <212> PRT  
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Pro Val Asp His Ile Ser Leu Ile Leu Ala Val Asp Trp Leu Val Asp  
 1 5 10 15

Arg Ser Cys Thr Val Leu Asn Val Glu Gly Asp Ala Leu Gly Ala Gly  
 20 25 30

Leu Leu Gln Asn Tyr Val Asp Arg Thr Glu Ser Arg Ser Thr Glu Pro  
 35 40 45

Glu Leu Ile Gln Val Lys Ser Glu Leu Pro Leu Asp Pro Leu Pro Val  
 50 55 60

Pro Thr Glu Glu Gly Asn Pro Leu Leu Lys His Tyr Arg Gly Pro Ala  
 65 70 75 80

Gly Asp Ala Thr Val Ala Ser Glu Lys Glu Ser Val Met  
 85 90

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Pro Thr Asp Asp Ile Thr Leu Ile Ile Ala Val Asp Trp Phe Leu Asp  
 1 5 10 15

Arg Leu Arg Thr Thr Thr Asn Val Leu Gly Asp Ser Leu Gly Ala Gly  
 20 25 30

Ile Val Glu His Leu Ser Arg His Glu Leu Lys Asn Arg Asp Val Glu  
 35 40 45

Met Gly Asn Ser Val Ile Glu Glu Asn Glu Met Lys Lys Pro Tyr Gln  
 50 55 60

Leu Ile Ala Gln Asp Asn Glu Thr Glu Lys Pro Ile Asp Ser Glu Thr  
 65 70 75 80

Lys Met

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 <212> PRT  
 <213> Rat P462-End rGlt-1b

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Pro Thr Glu Asp Ile Ser Leu Leu Val Ala Val Asp Trp Leu Leu Asp  
 1 5 10 15

Arg Met Arg Thr Ser Val Asn Val Val Gly Asp Ser Phe Gly Ala Gly  
 20 25 30

Ile Val Tyr His Leu Ser Lys Ser Glu Leu Asp Thr Ile Asp Ser Gln  
 35 40 45

His Arg Met His Glu Asp Ile Glu Met Thr Lys Thr Gln Ser Ile Tyr  
 50 55 60

Asp Asp Thr Lys Asn His Arg Glu Ser Asn Ser Asn Gln Cys Val Asn  
 65 70 75 80

Ala Ala His Asn Ser Val Val Ile Asp Glu Cys Lys Val Pro Phe Pro  
 85 90 95

Phe Leu Asp Ile Glu Thr Cys Ile  
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<211> 115  
 <212> PRT  
 <213> Rat P462-End rGLT-1

<400> 9

Pro Thr Glu Asp Ile Ser Leu Leu Val Ala Val Asp Trp Leu Leu Asp  
 1 5 10 15

Arg Met Arg Thr Ser Val Asn Val Val Gly Asp Ser Phe Gly Ala Gly  
 20 25 30

Ile Val Tyr His Leu Ser Lys Ser Glu Leu Asp Thr Ile Asp Ser Gln  
 35 40 45

His Arg Met His Glu Asp Ile Glu Met Thr Lys Thr Gln Ser Val Tyr  
 50 55 60

Asp Asp Thr Lys Asn His Arg Glu Ser Asn Ser Asn Gln Cys Val Tyr  
 65 70 75 80

Ala Ala His Asn Ser Val Val Ile Asp Glu Cys Lys Val Thr Leu Ala  
 85 90 95

Ala Asn Gly Lys Ser Ala Asp Cys Ser Val Glu Glu Glu Pro Trp Lys  
 100 105 110

Arg Glu Lys  
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 <213> Rat P486-End EAAT4

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Pro Thr Glu Asp Ile Thr Leu Ile Ile Ala Val Asp Trp Phe Leu Asp  
 1 5 10 15

Arg Leu Arg Thr Met Thr Asn Val Leu Gly Asp Ser Ile Gly Ala Ala  
 20 25 30

Val Ile Glu His Leu Ser Gln Arg Glu Leu Glu Leu Gln Glu Ala Glu  
 35 40 45

Leu Thr Leu Pro Ser Leu Gly Lys Pro Tyr Lys Ser Leu Met Ala Gln  
 50 55 60

Glu Lys Gly Ala Ser Arg Gly Arg Gly Gly Asn Glu Ser Ala Met  
65 70 75